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Effects of enzymatically hydrolyzed fish byproducts in diet of juvenile rainbow trout (*Oncorhynchus mykiss*)

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Abstract

Five experimental diets were formulated to evaluate the effects of dietary enzymatically hydrolyzed tuna by-product on growth, non-specific immune responses, and hematology of juvenile rainbow trout (*Oncorhynchus mykiss*). A basal diet with 50% of fishmeal was used as control (CON) and four other diets replaced 12.5% (TBB_{12.5}), 25% (TBB₂₅), 37.5% (TBB_{37.5}), and 50% (TBB₅₀) of fish meal in the CON diet. Juvenile rainbow trout (4.87 ± 0.05 g) were randomly distributed into 15 tanks (50 L) and fed 3–4% of wet body weight two times a day. At the end of 7 weeks of feeding trial, weight gain, specific growth rate, feed efficiency, and protein efficiency ratio of fish fed CON diet were significantly higher than those of fish fed TB₅₀ diet (P < 0.05). But there were no significant differences among fish fed CON, TBB_{12.5}, TBB₂₅, and TBB_{37.5} diets (P > 0.05). There were no significant differences in GPT levels among fish fed CON, TBB_{12.5}, TBB₂₅, and TBB_{37.5} diets. Also, there were no significant differences in lysozyme, superoxide dismutase, glucose, and total protein levels in all experimental diet (P > 0.05). The broken-line analysis indicated that the minimum dietary level of enzymatically hydrolyzed tuna by-product to replace fishmeal could be 29.7% in rainbow trout. These results indicated that the optimum level of dietary enzymatically hydrolyzed tuna by-product could replace greater than 29.7% but less than 37.5% of fishmeal in juvenile rainbow trout diet.

Keywords: Rainbow trout, Enzymatic hydrolysis, Growth performance, Immune response, Tuna by-products

Background

Fish meal (FM) is the one of the most important ingredients used in aquafeed. FM is a high-quality source of protein and is desirable for its rich essential amino acids profile (Cho and Kim 2011; Anderson et al. 2016) as well as for its vitamin and mineral content (IFOMA 2000). Typically, FM (66% crude protein) is the preferred source of protein in aquafeed formulation, yet the cost of FM is continually increasing (Tomas et al. 2005; Anderson et al. 2016; FAO 2018). It is to be noted that the world demand of FM increased by only about 27% during the past two decades, while FM output by the major FM-producing countries actually decreased. Because of the limited supply of FM coupled with its increasing demand, the cost of producing fish is likely to

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tuna muscle by-product powder is a promising feed ingredient to replace 50% FM protein without reduction in the growth performance of olive flounder.

Enzymatic hydrolysis is a process by which enzymes facilitate the cleavage of chemical bonds by water, playing a pivotal role in the digestion of food (Nielsang et al. 2005). This process makes it possible to improve the nutritive value of individual feed ingredients (PER, digestibility and FCR). Moreover, the moisture contents of the ingredients remain almost the same, varying only by 2.5 \pm 4.4%, maintaining typical parameters for storage of ingredients. Lastly, an important benefit of extrusion after enzymatic hydrolysis is the sterilization of the ingredients. Enzymatic hydrolyzed by-products are now widely used to modify or improve protein quality and their digestibility (Panyam and Kilara 1996; Liaset et al. 2000; Je et al. 2007; Baek and Cadwallader 1995; Nielsang et al. 2005). Nevertheless, the presence of endogenous enzymes in by-products may induce auto hydrolysis. In the case of squid gut and muscle, high endogenous proteolytic activity has been reported (Sugiyama et al. 1989; Leblanc and Gill 1982; Rodger et al. 1984) and during chopping, the tissues become hydrolyzed, rendering it richer in free essential amino acids (Lian et al. 2005). Hydrolyzed fish by-products from Alaskan fisheries have been evaluated for replacement of menhaden FM in Litopenaeus vannamei diets (Forster et al. 2011). Replacement of FM in aqua feeds with inexpensive protein sources, such as unexploited by-products which are regionally or locally available, may be a great opportunity for sustainable resource utilization.

Rainbow trout, Oncorhynchus mykiss, is one of most popular fish species around the world, due to its excellent meat quality (Vehviläinen et al. 2008). It is also one of the most studied species and is widely used for research in many fields such as carcinogenesis, toxicology, immunology, ecology, physiology, and nutrition (Thorgaard et al. 2002; Cowey 1992; Boyle et al. 2013; Berthelot et al. 2014). Global rainbow trout production from aquaculture has grown tremendously and has reached 814,090 metric tons as of 2016 (FAO 2018). For Republic of Korea, rainbow trout is also one of the commercially important freshwater species with 3358 metric tons of production in 2017; ranking third among freshwater species nationally (KOSTAT 2018). Numerous studies have been conducted on FM replacement in rainbow trout (Hauptman et al. 2014). However, to our knowledge, no study has been conducted using enzymatically hydrolyzed tuna by-products for replacing FM in the diet of juvenile rainbow trout. Based on the results obtained in previous studies, the aim of this study is to partially replace FM with enzymatically hydrolyzed tuna by-products in order to examine the effects on growth and non-specific immune responses in juvenile rainbow trout.

Materials and methods

Experimental design and diets

A basal diet was formulated containing crude protein 50.22%, crude lipid 13.71%, ash 9.79%, and moisture 4.93%; without inclusion of enzymatically hydrolyzed tuna backbone by-products, used as control diet (CON) and four other experimental diets were formulated using enzymatically hydrolyzed tuna backbone by-products. TBB_{12.5}, TBB₂₅, TBB_{37.5}, and TBB₅₀ diet were formulated with hydrolyzed tuna by-products level of 12.5%, 25%, 37.5%, and 50% replacement of FM respectively (Table 1). The ingredients were grounded into powder form using a grinding machine and sieved before inclusion of ingredients. The resultant mixtures become pelletized with a pellet machine with no heating using a module of 2-mm diameter and air dried for 3 days. The dried feeds were

Table 1 Formulation and proximate composition of five experimental diets (% of dry matter basis)

Ingredients	Diets ^a					
(% in diet)	CON	TBB _{12.5}	TBB ₂₅	TBB37.5	TBB ₅₀	
Fishmeal	50	43.75	37.50	31.25	25.00	
En. H.B ^b	0.00	5.82	11.64	17.46	23.27	
Blood meal	2.90	2.90	2.90	2.90	2.90	
Soybean meal	3.50	3.50	3.50	3.50	3.50	
Wheat gluten meal	7.9	7.9	7.9	7.9	7.9	
Cornstarch	0.00	0.6	1.18	1.78	2.39	
Wheat flour	20.00	20.00	20.00	20.00	20.00	
Soybean oil	7.80	7.80	7.80	7.80	7.80	
Fish oil	3.37	3.02	2.68	2.33	1.98	
Cellulose	0.53	0.71	0.90	1.08	1.26	
Vitamin premix	2.00	2.00	2.0	2.00	2.00	
Mineral premix	2.00	2.00	2.0	2.00	2.00	
Total	100	100	100	100	100	
Proximate analysis (% of DM)						
Moisture	4.93	5.72	6.17	7.22	7.30	
Protein	50.22	48.71	48.34	47.95	46.81	
Lipid	13.71	12.75	14.89	14.25	13.61	
Ash	9.79	9.59	9.49	9.11	9.05	

TBB12.5 tuna backbone by-products at 12.5% FM replacement (Samhae Company, Gyeongiu, South Korea)

TBB25 tuna backbone by-products at 25% FM replacement (Samhae Company, Gyeongiu, South Korea)

TBB37.5 tuna backbone by-products at 37.5% FM replacement (Samhae Company, Gyeongiu, South Korea)

TBB50 tuna backbone by-products at 50% FM replacement

^aCON control diet (the basal diet)

^bEn. H.B enzymatic hydrolyzed by-products (Samhae Company, Gyeongiu, South Korea)

sieved and packaged in zip-lock bags, labeled and stored at -20 °C until the feeding trial.

Experimental fish and feeding conditions

The feeding trial was conducted in the Department of Marine Bio-Materials and Aquaculture and Feeds and Foods Nutrition Centre, Pukyong National University (PKNU), Busan, Republic of Korea. Experimental rainbow trout fry were collected from a local rainbow trout farm named Ihwajeong Fish Farm, Sangju city, Korea. Before starting of the feeding trial, all fishes were fed with the basal diet two times in a day to acclimatize them to the laboratory experimental conditions for 2 weeks. Feeding trial was done using an indoor semi-recirculation system with 15 tanks of 50 L water holding capacity receiving constant filtered freshwater flow at a rate of 2 L min⁻¹ from the central tank. Half of the water was exchanged everyday by pumping filtered fresh water to the central tank. Supplementary continuous aeration facility provided to maintain required dissolved oxygen (D.O) and by using water heaters, temperature always maintained at 14.0 ± 0.1 °C throughout the experimental period. Four hundred eighty fishes with an average weight of 4.87 ± 0.05 g (mean \pm SD) were weighed and distributed as 20 fish/tank randomly into the 15 experimental indoor tanks. Three tanks were randomly selected for each one of the experimental diet. In this way, 15 tanks are randomly assigned for 5 dietary treatments and marked the tanks. Feeding was done two times daily (09:00 A.M. and 05:00 P.M.). Feed amount adjusted ranged from 3 to 4% according to the wet body weight in the beginning and at the end of feeding trial respectively for 7 weeks. Two hours after feeding, all tanks were siphoned to remove fecal matter and uneaten feeds daily. Mortality would check daily, dead fish immediately removed and weighed, and required feed amounts for the tanks adjusted according to the remaining body weight of fish in the tanks. Dead fish was not replaced during the experiment. Total weight of fish in each tank was determined at 4th week for adjust the amount of diet for feeding. Before each checking, fish were starved for 24 h to avoid inclusion of ingested feed in the measurements of fish body weight as well as to minimize stress.

Sample collection and analysis

After completion of the feeding trial, all fish were counted and weighed in each tank to calculate weight gain (WG), specific growth rate (SGR), feed efficiency (FE), protein efficiency ratio (PER), and survival rate (SR). From each tank, three fish were randomly selected, measurement of length, weight, and dissected to obtain visceral mass and liver to calculate hepatosomatic indices (HSI), viscerosomatic indices (VSI), and condition factor (CF). Blood samples were collected from the caudal vein of the fish with 1 ml syringes. Serum samples obtained from blood on clotting by centrifugation at $5000 \times g$ for 10 min and stored at – 70 °C for the analysis of the activities of lysozyme and superoxide dismutase (SOD). A turbid metric assay was used for the measurement of lysozyme activity of blood serum using the method by Hultmark (1980) with slightly modifications. Micrococcus lysodeikticus (0.75 mg ml⁻¹) was suspended in sodium phosphate buffer (0.1 M, pH 6.4), then 200 μ l of suspension was placed in each well of 96-well plates and after that 20 µl of test serum was added. After incubation, the reduction in absorbance of the samples was recorded at 570 nm at room temperature for 0 and 60 min in a micro plate reader (UVM 340, Biochrom, Cambridge, UK). One unit of lysozyme activity was regarded as a reduction in absorbance of 0.001 min^{-1} .

Activity of SOD was measured according to the percentage of reaction inhibition rate of the enzyme with WST-1 (water-soluble tetrazolium dye) substrate and xanthine oxidase using an SOD Assay Kit (Enzo ADI-900-157, Enzo Life Sciences, Inc.) in accordance with manufacturer's instructions. Each endpoint assay was monitored by absorbance at 450 nm (the absorbance wavelength for the colored product of WST-1 reaction with superoxide) after 20 min of reaction time at 37 °C. The inhibition percentage was normalized by mg of protein and showed as SOD activity units. Serum portions were also used for the biochemical parameters analysis of total protein (TP) of plasma, glutamic oxaloacetic transaminase (GOT), glutamic pyruvic transaminase (GPT), and glucose activities determined using the commercial clinical kits of Fuji DRI-CHEM 3500i, Fuji Photo Film Ltd., Tokyo, Japan.

From each tank, three fish were collected and pooled together according to the fish fed with different experimental diets; homogenized and freeze-dried at - 80 °C for analysis. Proximate compositions of the experimental fish and the basal diet were determined according to standard procedures of AOAC (2005) in duplicates. Both samples were dried at 105 °C to a constant weight to determine moisture content. Determination of ash content was done by incineration in a muffle furnace at 550 °C for 3.5 h, crude lipid was determined by the Soxhlet extraction using Soxtec system 1046 (Tecator AB, Foss, Hoganas, Sweden) and crude protein by the Kjedahl method (N \times 6.25) after acid digestion. Laboratory analyses were done at the Feeds and Foods Nutrition Research Centre (FFNRC) at Yongdang campus of PKNU, Republic of Korea.

Statistical analysis

After completion of all laboratory analysis, all data were analyzed by the one-way analysis of variance (ANOVA) using SAS Program, Version 9.1 (SAS Institute, Cary, NC, USA) to test for the effects of the different dietary experimental treatments. A least significant difference (LSD) test was used to compare the means of significant effect. Effects of treatments were considered significant at confidence level of P < 0.05. For each treatment, mean and SD were also calculated.

Results

Growth performances

Growth performances of juvenile rainbow trout, Oncorhynchus mykiss, based on weight gain (WG), specific growth rate (SGR), feed efficiency (FE), protein efficiency ratio (PER), and survival rate (SR) fed with five different experimental diets for 7 weeks are summarized in Table 2. After completion of feeding trial, WG, SGR, FE, and PER of fish fed CON was significantly higher than fish fed TBB₅₀ diet (P < 0.05). But there were no significant differences (P > 0.05) between fish fed TBB_{12.5}, TBB₂₅, and TBB_{37.5} diets. Broken-line model analysis on the basis of WG indicated that the minimum dietary level of enzymatic hydrolyzed of tuna backbone by-product to replace fishmeal could be up to 29.7% in rainbow trout (Fig. 1). Survival rate varied from 91.67 to 95.67%, but no significant difference (P > 0.05) was found among fish fed with all dietary treatments. There was no significant difference in hepatosomatic index (HSI). Viscerosomatic index (VSI) of fish fed TBB_{37.5} was significantly higher than fish fed TBB₅₀ diet. But no significant difference was found among the fish fed CON, TBB_{12.5}, and TBB₂₅ diets. CF of CON was significantly higher than TBB₁₂ and TBB₅₀ diets. But no significant difference was found among the fish fed TBB₂₅ and TBB_{37.5} diets.

Non-specific immune responses

Non-specific immune responses based on lysozyme and superoxide dismutase (SOD) activities of juvenile rainbow trout fed with five different experimental diets for 7 weeks are summarized in Table 3. In lysozyme activity and superoxide dismutase, no significant differences were found among the fish fed every experimental groups (P < 0.05).

Whole body proximate analysis

Whole body proximate compositions of juvenile rainbow trout fed with five different experimental diets for 7 weeks are summarized in Table 4. Moisture of fish fed TBB_{37.5} diet was significantly higher than CON, TBB_{12.5}, TBB₂₅, and TBB₅₀ diets. No significant differences were found among the fish fed CON, TBB_{12.5}, TBB₂₅, and TBB₅₀ diets. Crude protein of fish fed CON showed no significant differences among the fish fed every diets. But fish fed TBB_{12.5}, TBB₂₅, and TBB₅₀ diets were significantly higher than fish fed TBB_{37.5} diet. No significant difference was founded in crude lipid and crude ash among the fish fed every diets.

Hematological parameters

Hematological parameters of blood serum in juvenile rainbow trout fed with five different experimental diets for 7 weeks are summarized in Table 5. Total protein of fish fed CON showed no significant differences (P > 0.05) among the fish fed all the experimental diets. But fish fed TBB_{12.5} diet was significantly higher (P < 0.05) than TBB_{37.5} and TBB₅₀ diets. GOT of fish fed TBB₂₅, TBB_{37.5}, and TBB₅₀ diets was significantly higher (P < 0.05) than fish fed CON diet. But fish fed CON and TBB_{12.5} diets showed no significant differences

Table 2 Growth performances of juvenile rainbow trout fed with five experimental diets for 7 weeks

	Diets					
	CON	TBB _{12.5}	TBB ₂₅	TBB _{37.5}	TBB ₅₀	
WG ¹	259 ± 14.9^{a}	250 ± 17.4^{ab}	243 ± 17.7^{ab}	239 ± 29.0^{ab}	225 ± 8.82^{b}	
SGR ²	2.13 ± 0.07^{a}	2.09 ± 0.08^{ab}	2.05 ± 0.09^{ab}	2.03 ± 0.14^{ab}	1.96 ± 0.05^{b}	
FE ³	133 ± 6.14^{a}	131 ± 6.34^{ab}	127 ± 1.30^{ab}	125 ± 2.82^{ab}	$125\pm1.88^{\rm b}$	
PER^4	2.76 ± 0.12^{a}	2.70 ± 0.12^{ab}	2.63 ± 0.03^{ab}	2.62 ± 0.05^{ab}	2.59 ± 0.04^{b}	
HSI ⁵	1.07 ± 0.50^{ns}	0.91 ± 0.15	0.98 ± 0.12	1.03 ± 0.15	1.00 ± 0.13	
VSI ⁶	6.58 ± 0.67^{ab}	7.01 ± 0.60^{ab}	6.87 ± 1.13^{ab}	$7.26\pm0.99^{\rm a}$	6.35 ± 0.53^{b}	
CF ⁷	$1.07\pm0.08^{\rm a}$	0.98 ± 0.09^{bc}	$1.04\pm0.08^{\rm ab}$	0.98 ± 0.06^{ab}	$0.90 \pm 0.06^{\circ}$	
SR ⁸	93.3 ± 7.64^{ns}	91.7 ± 5.77	91.7 ± 7.64	93.3 ± 11.6	95.7 ± 2.89	

Values are means from triplicate groups of fish (n = 3) where values in each row with different superscripts are significantly different (P < 0.05)

¹Weight gain (WG) = (final weight – initial weight) \times 100/initial weight

²Specific growth rate (SGR; %/day) = (In final weight – In initial weight) × 100/d

³Feed efficiency (FE; %) = wet WG (g) \times 100/dry feed intake (g)

⁴Protein efficiency ratio (PER) = wet weight gain/protein intake

⁵Hepatosomatic index (HSI; %) = (liver weight/body weight) × 100

⁶Viscerosomatic index (VSI; %) = (visceral weight/body weight) \times 100

⁷Condition factor (CF) = {fish weight (g)/fish length (cm) 3 × 100

⁸Survival (SR; %) = (total fish-dead fish) \times 100/total fish



(P > 0.05). GPT of fish fed TBB₅₀ was significantly higher (P < 0.05) than the other diets. But no significant difference (P > 0.05) was observed among fish fed CON, TBB_{12.5}, TBB₂₅, TBB_{37.5} and TBB₅₀.

Discussion

In this study, enzymatically hydrolyzed tuna backbone by-product was used in different levels in experimental diets to partially replace commercially used fish meal (FM). After completion of the feeding trial, weight gain (WG) and specific growth rate (SGR) of fish fed the 37.5% replacement diets with tuna backbone by-product were not significantly different from the FM fed fish. Although, 50% replacement of FM with tuna backbone by-product reduced both the WG and SGR. This indicates that enzymatic hydrolyzed tuna backbone by-product containing diets can partially replace FM but up to at a certain level. In a relevant study by Mendoza et al. (2001), it was reported that using enzymatic hydrolyzed and co-extruded feather meal with soya-bean meal in practical diets for the pacific-white shrimp (Litopenaeus vannamei) can replace up to 20% of FM without

Table 3 Non-specific immune responses of juvenile rainbow troutfed with five experimental diets for 7 weeks

	Diets					
	CON	TBB ₁₂	TBB ₂₅	TBB37	TBB ₅₀	
Lys ¹	$1.35\pm0.30^{\text{ns}}$	1.58 ± 0.32	1.60 ± 0.28	1.64 ± 0.37	1.24 ± 0.05	
SOD ²	$80.9\pm8.27^{\text{ns}}$	79.0 ± 7.76	80.6 ± 4.09	76.4 ± 2.02	83.6 ± 7.07	

Values are means from triplicate groups of fish (n = 3) where in each row with different superscripts values are significantly different (P < 0.05) ¹Lysozyme (U/I)

²Superoxide dismutase (% inhibition)

reducing growth performance and body composition. Feed utilization performance parameters such as FE and PER were also influenced by dietary treatments in this study. FE and PER were both similar in fish fed the CON diet and those of fish fed diets replacing up to 37.5% of fish meal with tuna backbone by-product. Higher replacement levels (TBB₅₀) decreased FE and PER in rainbow trout. This indicates that up to 37.5% of FM replacement by tuna backbone by-product did not negatively influence feed utilization performance of rainbow trout. These results were consistent with the observations in studies that used combinations of various protein sources in the diets of rainbow trout (Jo et al. 2016; Bureau et al. 2000). Similar studies on rainbow trout have also reported that partially hydrolyzed ingredients have higher nutritional value because of higher

Table 4 Whole body proximate composition of juvenilerainbow trout fed with five experimental diets for 7 weeks(% dry matter basis)

	Diets					
	CON	TBB _{12.5}	TBB ₂₅	TBB _{37.5}	TBB ₅₀	
Mo ¹	78.3 ± 0.11^{b}	77.2 ± 0.58^{b}	78.3 ± 0.22^{b}	79.1 ± 0.32^{a}	78.3 ± 0.47 ^b	
CP^2	14.6 ± 0.45^{ab}	15.1 ± 0.25^{a}	15.1 ± 0.02^{a}	$14.4\pm0.07^{\rm b}$	15.2 ± 0.56^{a}	
CL ³	$4.80\pm0.75^{\text{ns}}$	5.06 ± 0.52	5.26 ± 0.06	4.60 ± 0.20	4.93 ± 0.44	
As^4	$1.94\pm0.10^{\text{ns}}$	2.06 ± 0.10	1.96 ± 0.10	1.92 ± 0.05	2.10 ± 0.16	

Values are means from triplicate groups of fish (n = 3) where in each row with different superscripts values are significantly different (P < 0.05)

¹*Mo* moisture content of whole body composition of juvenile rainbow trout in different diets

²CP crude protein present under different dietary treatments

³*CL* crude lipid, present in whole body proximate composition of juvenile rainbow trout after feeding trial

⁴As ash content of whole body proximate composition of fish in different diets

	Diets					
	CONT	TBB _{12.5}	TBB ₂₅	TBB _{37.5}	TBB ₅₀	
GL ¹	105 ± 6.51^{ns}	112±8.19	113±3.00	111.3 ± 6.03	107 ± 9.17	
TP ²	3.60 ± 0.10^{ab}	3.73 ± 0.23^{a}	$3.40\pm0.26^{\text{ab}}$	3.30 ± 0.10^{b}	3.33 ± 0.25^{b}	
GOT ³	$402 \pm 39.1^{\circ}$	452 ± 26.0^{bc}	517 ± 16.2^{ab}	530 ± 52.9^{a}	509 ± 44.5^{ab}	
GPT ⁴	8.00 ± 3.61^{b}	8.67 ± 2.08^{b}	10.3 ± 2.31^{b}	10.0 ± 1.73^{b}	18.7 ± 2.52^{a}	

Table 5 Hematological parameters of juvenile rainbow trout fed with five experimental diets for 7 weeks

Values are means from triplicate groups of fish (n = 3) where in each row with different superscripts values are significantly different (P < 0.05) ¹GL glucose (mg/dl)

²TP total protein (g/dl)

³GOT glutamic glutamic oxaloacetic transaminase (U/I)

⁴GPT glutamic pyruvic transaminase (U/I):

levels of free amino acids. This makes these ingredients more digestible and fish can utilize the ingredients more efficiently that will results in higher growth and feed utilization performances (Dong et al. 1993; Hardy 2002). The hepatosomatic index (HSI) was not changed in fish fed all the experimental diets. Although, viscerosomatic index (VSI) and condition factor (CF) were lower in fish fed the TBB50 diet compared to those of fish fed the CON diet. HIS, VSI, and CF are good representatives for health, general well-being, and feed quality (Ighwela et al. 2014). These parameter showed that until 37.5% replacement of FM with tuna backbone by-product has no negative effects on rainbow trout organosomatic health. These results were similar with previous studies using tuna by-products as a protein source for rainbow trout (Samuelsen et al. 2001; Caballero et al. 2002; Tekinay et al. 2009). Noteworthy, in this study, we observed more than 90% survival for all experimental groups that could be related to standard rearing condition and feed. Whole body proximate composition of rainbow trout fed the experimental diets did not show any specific trend in the present study. Whereas crude lipid and ash were same for all fish fed the experimental diets and there were no significant differences in crude protein between the CON and other experimental groups. Some studies have reported a trend in which whole body moisture content shows opposite values compared to protein content in rainbow trout, O. mykiss (Jo et al. 2016); kuruma shrimp, *Marsupenaeus japonicas* (Bulbul et al. 2016); and Australian short-finned eel, Aaustralis australis (Engin and Carter 2005).

Non-specific immune responses are the basic mechanism of defense system in fish and play a vital role in the obtained immune response and homeostasis by a system of receptor proteins (Srivastava and Pandey 2015; Halver and Hardy 2002; Reverter et al. 2014; Uribe et al. 2011). Lysozyme activity is an important humoral indicator of non-specific immunity in fish and one of the important enzymes of defensive mechanism that protects the host (Ellis 1999; Jo et al. 2016). On the other hand, SOD is a metalloenzyme that catalyzes the superoxide radical (O_{2-}) dismutation into molecular oxygen (O_2) and hydrogen peroxide (H_2O_2) (Shao et al. 2010). This will support a significant defense system against oxidative damage. Findings of the present study did not show any significant differences in lysozyme or SOD activities in all the experimental fish. This shows that even 50% replacement of FM with tuna backbone by-product does not influence the innate immunity of rainbow trout. This could be attributed to standard experimental conditions while tuna backbone by-product was providing sufficient protein with well-balanced amino acids to support the immune system (Sullivan 2008; Jo et al. 2016).

Serum parameters as well as hematological activities are good indicators of health status of any organism and can vary with temperature, season, and nutritional status (Blaxhall 1972). They are suitable indicators for evaluating the health, stress level, and physiological welfare of fish (Simide et al. 2016). Serum glucose is a stress indicator of fish and stress hormone which is associated with cortisol, mobilize and increase glucose production in blood serum to cope with the required energy produced by a stressor (Wallace et al. 1994). Total protein of blood serum is also considered a strong natural response in fish (Reverter et al. 2014) and assessments can reflect nutritional levels. (Acharya and Mohanty 2014). In the present study, glucose content of serum did not show any significant differences in fish fed all the experimental diets. There were also no differences in serum total protein level of fish fed the CON diet and those of fish fed the other diets. Glutamic oxaloacetic transaminase (GOT) and glutamic pyruvic transaminase (GPT) which are enzymes or proteins mostly found in liver cells and rising levels in blood serum indicate damage or inflammation of liver cells (Giannini et al. 2005). In the present study, the GOT level showed an increasing trend with higher FM replacement levels. While fish fed diets TBB₂₅, TBB_{37.5}, and TBB₅₀ had a higher GOT compared to fish fed the CON diet. Serum GPT also showed higher levels for fish fed the TBB₅₀ diet compared to all other experimental fish. An increase in GOT and GPT could be attributed to the addition of alternative protein

sources in fish diets because of various dynamics such as reduction in FM content, tolerance of anti-nutritional factors, digestibility, and palatability problems (Jo et al. 2016; National Research Council, NRC 2011).

Conclusions

Considering growth performances, non-specific immune responses, other relevant parameters in CON, and other diets which included tuna backbone by-products for FM replacement, the present study demonstrated that FM can be replaced greater than 29.7% but less than 37.5% by enzymatically hydrolyzed tuna backbone by-products in the diets of juvenile rainbow trout. The findings of the study will also encourage the feed manufacturers to utilize enzymatically hydrolyzed tuna backbone by-products as alternative protein sources in generating low-cost and healthy aqua feed.

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Availability of data and materials

Please contact author for data requests.

Authors' contributions

JB designed, conducted the experiment, and wrote the paper. MDAKA conducted feeding trial. SW design and conducted some parts of the analysis and experiment. AH prepared of the ingredients and feeding trial. MS conducted analysis of sample. SCB supervised the experiment. All authors read and approved the final manuscript.

Ethics approval and consent to participate

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Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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